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SEQUENCE LISTING

<110> CHUGAI SEIYAKU KABUSHIKI KAISHA

<120> Expression systems using mammalian β -Actin promoter

<130> C1-A0311P

<150> JP 2003-405269

<151> 2003-12-03

<160> 39

<170> PatentIn version 3.1

<210> 1

<211> 1577

<212> DNA

<213> Mus musculus

<400> 1

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1577

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 <213> Woodchuck hepatitis virus

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 <212> DNA
 <213> Homo sapiens

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<211> 660

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<213> Mus musculus

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tctactggac atcttagaca cagcaggtca agaagagtat agtgccatgc gggaccagta 240

catgcgcaca ggggagggct tcctctgtgt atttgccatc aacaacacca agtccttcga 300

ggacatccat cagtacaggg agcagatcaa gcggtgaaa gattcagatg atgtgccaat 360

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gaacccaccc gatgagagtg gtcctggctg catgagctgc aaatgtgtgc tgcctgaca 600

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<211> 576

<212> DNA

<213> Mus musculus

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ctctgttat ttgccatcaa caacaccaag tccttcgagg acatccatca gtacagggag 300

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<211> 189

<212> PRT

<213> Mus musculus

<400> 7

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Asp Pro Thr Ile Glu Asp Ser Tyr Arg Lys Gln Val Val Ile Asp Gly
 35 40 45

Glu Thr Cys Leu Leu Asp Ile Leu Asp Thr Ala Gly Gln Glu Glu Tyr
 50 55 60

Ser Ala Met Arg Asp Gln Tyr Met Arg Thr Gly Glu Gly Phe Leu Cys
 65 70 75 80

Val Phe Ala Ile Asn Asn Thr Lys Ser Phe Glu Asp Ile His Gln Tyr
 85 90 95

Arg Glu Gln Ile Lys Arg Val Lys Asp Ser Asp Asp Val Pro Met Val
 100 105 110

Leu Val Gly Asn Lys Cys Asp Leu Ala Ala Arg Thr Val Glu Ser Arg
 115 120 125

Gln Ala Gln Asp Leu Ala Arg Ser Tyr Gly Ile Pro Tyr Ile Glu Thr
 130 135 140

Ser Ala Lys Thr Arg Gln Gly Val Glu Asp Ala Phe Tyr Thr Leu Val
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 165 170 175

Ser Gly Pro Gly Cys Met Ser Cys Lys Cys Val Leu Ser
 180 185

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<211> 188

<212> PRT

<213> Homo sapiens

<400> 8

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1 5 10 15

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20 25 30

Asp Pro Thr Ile Glu Asp Ser Tyr Arg Lys Gin Val Val Ile Asp Gly

35 40 45

Glu Thr Cys Leu Leu Asp Ile Leu Asp Thr Ala Gly Gin Glu Glu Tyr

50 55 60

Ser Ala Met Arg Asp Gin Tyr Met Arg Thr Gly Glu Gly Phe Leu Cys

65 70 75 80

Val Phe Ala Ile Asn Asn Thr Lys Ser Phe Glu Asp Ile His His Tyr

85 90 95

Arg Glu Gin Ile Lys Arg Val Lys Asp Ser Glu Asp Val Pro Met Val

100 105 110

Leu Val Gly Asn Lys Cys Asp Leu Pro Ser Arg Thr Val Asp Thr Lys

115 120 125

Gin Ala Gin Asp Leu Ala Arg Ser Tyr Gly Ile Pro Phe Ile Glu Thr

130 135 140

Ser Ala Lys Thr Arg Gin Gly Val Asp Asp Ala Phe Tyr Thr Leu Val

145 150 155 160

Arg Glu Ile Arg Lys His Lys Glu Lys Met Ser Lys Asp Gly Lys Lys

165 170 175

Lys Lys Lys Ser Lys Thr Lys Cys Val Ile Met

180

185

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<211> 27

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Pri
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27

<210> 10

<211> 27

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Pri
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27

<210> 11

<211> 26

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Pri

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26

<210> 12

<211> 26

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Pri
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<400> 12

aagcttggcg aactatcaag acacaa

26

<210> 13

<211> 50

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Pri
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<400> 13

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<210> 14

<211> 50

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Pri
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<400> 14

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<210> 15

<211> 50

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Pri
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<210> 16

<211> 50

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Pri
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<400> 16

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<210> 17

<211> 50

<212> DNA

<213> Artificial

<220>

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<400> 17

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<210> 18

<211> 50

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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<210> 19

<211> 50

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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<210> 20

<211> 50

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 20

gtcccgaaaa ggagctgaca ggtggggca atgccccaaac cagtgggggt

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<210> 21

<211> 50

<212> DNA

<213> Artificial

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50

<210> 22

<211> 50

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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<210> 23

<211> 50

<212> DNA

<213> Artificial

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<211> 50

<212> DNA

<213> Artificial

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<210> 25

<211> 50

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 25

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<210> 26

<211> 50

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 26

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<210> 27

<211> 50

<212> DNA

<213> Artificial

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<210> 28

<211> 57

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 28

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<210> 29

<211> 56

<212> DNA

<213> Artificial

<220>

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<210> 30

<211> 30

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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30

<210> 31

<211> 26

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 31

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<210> 32

<211> 25

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 32

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25

<210> 33

<211> 22

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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<210> 34

<211> 27

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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27

<210> 35

<211> 27

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 35

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27

<210> 36

<211> 24

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

<400> 36

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24

<210> 37

<211> 20

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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<210> 38

<211> 27

<212> DNA

<213> Artificial

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<223> Description of Artificial Sequence : Artificially Synthesized Primer Sequence

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27

<210> 39

<211> 27

<212> DNA

<213> Artificial

<220>

<223> Description of Artificial Sequence : Artificially Synthesized Primary Sequence

<400> 39

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27